

[illegible][illegible]

NTDB id 587	KW2 RS10850 WP 042211510.1	GALIVLLTSLSIIIIFKHHISALTNFTFLVKVPLVSSFIRLYLTAYFSREWGNLIAQGVVELRQIIELMKKKQKSRIFAEVGIKLDLGL	251
NTDB id 82977	H7678 RS00730 WP 050807598.1	ILFIGVLLGGIWLHWRSSQRLKLFTHFSAYPVLGKLLQQYLTAYYAREWGTLIGQGLELPVILDIMAMEKSLFMQELAEIDIRVSL	253
NTDB id 395	FSA28 RS09400 WP 257638827.1	CCFFILLVGIIFTTFSSRKISRHLIGFCSHLPFVGKYLQLYLTAYYAREWGNLIGQGVEMAQIVRLMQDQKSQQLFREICYDMEKAL	253
NTDB id 386	SMU RS09030 WP 255262721.1	CCFFILLVGIIFTTFSSRKISRHLIGFCSHLPFAGKYLQLYLTAYYAREWGNLIGQGVEMAQIVRLMQDQKSQQLFREICYDMEKAL	253
NTDB id 430	SGO RS09415 WP 012130941.1	VLMLLGLLTGAIYLVFKGQKRPVYSFLARLPFVGSFVRIYLTAYYAREWGNMIGQGLELSQIFQIMQEQRSVLFQEQIGQDLGQAL	254
NTDB id 271	KZH43 RS09260 WP 074196785.1	VGLVSVLALLALTIFYKRSKMSVFSILARLPFIGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMQEQQSQLFKEVGQDLAQTL	248
NTDB id 230	SPD RS09890 WP 074196785.1	VGLVSVLALLALTIFYKRSKMSVFSILARLPFIGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMQEQQSQLFKEVGQDLAQTL	248
NTDB id 196	SPR RS09425 WP 074196785.1	VGLVSVLALLALTIFYKRSKMSVFSILARLPFIGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMQEQQSQLFKEVGQDLAQTL	248
NTDB id 161	SP RS10375 WP 074196785.1	VGLVSVLALLALTIFYKRSKMSVFSILARLPFIGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMQEQQSQLFKEVGQDLAQTL	248
NTDB id 529	SMSK321 RS09235 WP 080550783.1	VGLVSLTLLALTIFYKRSKMRVFSILARIPFLGIFVQTYLTAYYAREWGNMISQGMELTQIFQIMQEQQSQLFKEIGQDLAQSL	248
NTDB id 501	SM12261 RS00950 WP 004238896.1	VGFSVSGVLLALTIFYKRGSKMRVFSILARIPFLGIFVQTYLTAYYAREWGNMISQGMELTQIFQIMQEQQSQLFKEIGQDMAHAL	248
consensus		***** * ***** * * * * * ! * * * * * ! ! ! ! * ! ! ! ! * ! * ! * ! * ! * * ! * * * ! * * * ! * * * * * !	

